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=> s dna probe and expansin activity
 L1 0 DNA PROBE AND EXPANSIN ACTIVITY

=> s dna probe and expansin
 L2 0 DNA PROBE AND EXPANSIN

=> s dna and probe and expansin
 L3 1 DNA AND PROBE AND EXPANSIN

=> d l3 ibib ab

L3 ANSWER 1 OF 1 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.
 ACCESSION NUMBER: 2001:277733 BIOSIS
 DOCUMENT NUMBER: PREV200100277733
 TITLE: Differential RNA expression of alpha-**expansin**
 gene family members in the parasitic angiosperm *Triphysaria*
versicolor (Scrophulariaceae).
 AUTHOR(S): Wrobel, Russell L.; Yoder, John I. (1)
 CORPORATE SOURCE: (1) Department of Vegetable Crops, University of
 California, Davis, 1 Shields Ave., Davis, CA, 95616:
 jiyoder@ucdavis.edu USA
 SOURCE: Gene (Amsterdam), (21 March, 2001) Vol. 266, No. 1-2, pp.
 85-93. print.
 ISSN: 0378-1119.
 DOCUMENT TYPE: Article
 LANGUAGE: English
 SUMMARY LANGUAGE: English

AB Haustoria are parasitic plant specific organs that locate, attach to, and invade host plant tissues. Parasitic species of the Scrophulariaceae develop haustoria on their roots in response to chemical signals released by host plant roots. Haustorium development was induced in vitro in roots of the parasitic Scrophulariaceae *Triphysaria versicolor* by treating them with exudates obtained from maize roots, the chemical 2,6-dimethoxybenzoquinone (DMBQ) or the cytokinin 6-benzylaminopurine (BAP). Morphological responses of *T. versicolor* roots to these haustoria inducing factors (HIFs) included localized swelling and epidermal hair proliferation near the root tips. These responses were not observed when roots of the non-parasitic Scrophulariaceae *Lindenbergia muraria* were similarly treated. Because **expansin** proteins are closely

associated with plant cell wall expansion and growth, we examined the expression of **expansin** genes in response to HIFs. We isolated cDNAs homologous to transcripts encoding three distinct alpha-**expansin** proteins in *T. versicolor*. Northern-blot analyses indicated that these transcripts were differentially abundant in different tissues. Steady-state levels of two **expansin** transcripts increased in *T. versicolor* roots exposed to BAP, but not DMBQ or maize root exudates. **Expansin** transcript abundance also increased in *L. muraria* in response to BAP treatment. These results suggest that the expansins examined fulfill functions distinct from haustorium development.

=> s expansin protein and dna

L4 4 EXPANSIN PROTEIN AND DNA

=> dup rem l4

PROCESSING COMPLETED FOR L4

L5 4 DUP REM L4 (0 DUPLICATES REMOVED)

=> s l5 and probe

L6 0 L5 AND PROBE

=> d l5 1-4 ibib ab

L5 ANSWER 1 OF 4 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2002:754532 CAPLUS

DOCUMENT NUMBER: 137:274419

TITLE: Protein and cDNA sequences of .beta.-**expansin**

protein isolated from maize and polynucleotides and methods of uses thereof

INVENTOR(S): Multani, Dilbag S.; Johal, Gurmukh S.

PATENT ASSIGNEE(S): Pioneer Hi-Bred International, Inc., USA

SOURCE: PCT Int. Appl., 65 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2002077170	A2	20021003	WO 2002-US8603	20020320
W:	AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG			

PRIORITY APPLN. INFO.: US 2001-277847P P 20010322

US 2001-324182P P 20010921

AB The present invention provides protein and cDNA sequences of .beta.-**expansin protein** isolated from maize and methods for modulating plant cell enlargement, plant strength, plant pliability and flexibility. Specifically, the invention discloses that the sequence can be used in expression cassettes for modulating plant cell enlargement, stalk strength, plant pliability and flexibility. Transformed plants, plant cells, tissues, and seed are also provided. Methods for rapidly identifying and isolating a Mu-tagged recessive gene mutation in a F1 generation plant, and identification and isolation of its assocd. wild-type gene are also provided.

L5 ANSWER 2 OF 4 MEDLINE
 ACCESSION NUMBER: 2001406338 MEDLINE
 DOCUMENT NUMBER: 21351003 PubMed ID: 11457903
 TITLE: Expression of six expansin genes in relation to extension activity in developing strawberry fruit.
 AUTHOR: Harrison E P; McQueen-Mason S J; Manning K
 CORPORATE SOURCE: Horticulture Research International, Wellesbourne, Warwick CV35 9EF, UK.. elizabeth.harrison@hri.ac.uk
 SOURCE: JOURNAL OF EXPERIMENTAL BOTANY, (2001 Jul) 52 (360) 1437-46.
 Journal code: 9882906. ISSN: 0022-0957.
 PUB. COUNTRY: England: United Kingdom
 DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
 LANGUAGE: English
 FILE SEGMENT: Priority Journals
 OTHER SOURCE: GENBANK-AF226700; GENBANK-AF226701; GENBANK-AF226702; GENBANK-AF226703; GENBANK-AF226704
 ENTRY MONTH: 200110
 ENTRY DATE: Entered STN: 20011008
 Last Updated on STN: 20011008
 Entered Medline: 20011004

AB Expansins are proteins which have been demonstrated to induce cell wall extension in vitro. The identification and characterization of six expansin cDNAs from strawberry fruit, termed FaExp3 to FaExp7, as well as the previously identified FaExp2 is reported here. Analysis of expansin mRNAs during fruit development and in leaves, roots and stolons revealed a unique pattern of expression for each cDNA. FaExp3 mRNA was present at much lower levels than the other expansin mRNAs and was expressed in small green fruit and in ripe fruit. FaExp4 mRNA was present throughout fruit development, but was more strongly expressed during ripening. FaExp5 was the only clone to show fruit specific expression which was up-regulated at the onset of ripening. FaExp6 and FaExp7 mRNAs were present at low levels in the fruit with highest expression in stolon tissue. During fruit development FaExp6 had the highest expression at the white, turning and orange stages whereas expression of FaExp7 was highest in white fruit. The expression profiles of FaExp2 and FaExp5 in developing fruit were similar except that FaExp2 was induced at an earlier stage. Analysis of **expansin protein** by Western blotting using an antibody raised against CsExp1 from cucumber hypocotyls identified two bands of 29 and 31 kDa from developing fruit. Protein extracts from developing fruit were assayed for extension activity. Considerable rates of extension were observed with extracts from ripening fruit, but no extension was observed with protein from unripe green fruit. These results demonstrate the presence of at least six expansin genes in strawberry fruit and that during ripening the fruit acquires the ability to cause extension in vitro, characteristic of expansin action.

L5 ANSWER 3 OF 4 MEDLINE
 ACCESSION NUMBER: 1998393519 MEDLINE
 DOCUMENT NUMBER: 98393519 PubMed ID: 9724690
 TITLE: Localized upregulation of a new expansin gene predicts the site of leaf formation in the tomato meristem.
 AUTHOR: Reinhardt D; Wittwer F; Mandel T; Kuhlmeier C
 CORPORATE SOURCE: Institute of Plant Physiology, University of Berne, Altenbergrain 21, CH-3013 Berne, Switzerland.
 SOURCE: PLANT CELL, (1998 Sep) 10 (9) 1427-37.
 Journal code: 9208688. ISSN: 1040-4651.
 PUB. COUNTRY: United States
 DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
 LANGUAGE: English
 FILE SEGMENT: Priority Journals
 ENTRY MONTH: 199811

ENTRY DATE: Entered STN: 19990106
Last Updated on STN: 19990106
Entered Medline: 19981118

AB Expansins are extracellular proteins that increase plant cell wall extensibility in vitro and are thought to be involved in cell expansion. We showed in a previous study that administration of an exogenous **expansin protein** can trigger the initiation of leaflike structures on the shoot apical meristem of tomato. Here, we studied the expression patterns of two tomato expansin genes, LeExp2 and LeExp18. LeExp2 is preferentially expressed in expanding tissues, whereas LeExp18 is expressed preferentially in tissues with meristematic activity. In situ hybridization experiments showed that LeExp18 expression is elevated in a group of cells, called I1, which is the site of incipient leaf primordium initiation. Thus, LeExp18 expression is a molecular marker for leaf initiation, predicting the site of primordium formation at a time before histological changes can be detected. We propose a model for the regulation of phyllotaxis that postulates a crucial role for expansin in leaf primordium initiation.

L5 ANSWER 4 OF 4 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1997:34082 CAPLUS
DOCUMENT NUMBER: 126:56632
TITLE: Purified expansin proteins and their effects on cellulose paper
INVENTOR(S): Cosgrove, Daniel J.; Mcqueen-Mason, Simon; Gultinan, Mark; Shcherban, Tatyana; Shi, Jun
PATENT ASSIGNEE(S): Penn State Research Foundation, USA
SOURCE: PCT Int. Appl., 92 pp.
CODEN: PIXXD2
DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 5
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9635442	A1	19961114	WO 1996-US6759	19960513
W: CA, JP, KR				
RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
US 5959082	A	19990928	US 1995-440517	19950512
PRIORITY APPLN. INFO.: US 1995-440517 A 19950512				
US 1993-60944 B2 19930512				
US 1994-242090 B2 19940512				

AB The present invention relates to a new class of proteins, known as expansins, and their isolation, sequencing, genesis by expression systems, and utilization. Thus, the walls of growing cucumber seedlings possess extractable proteins which can induce extension of isolated walls. The names expansin-29 and expansin-30 were proposed for the 2 specific members of this class, based on their relative mol. masses on SDS-PAGE. Three peptide fragments from the purified cucumber Ex-29 protein were sequenced, oligonucleotide primers designed to amplify a portion of the expansin cDNA using PCR, and the PCR fragment used to screen a cDNA library to identify full-length clones. Expansin proteins were also purified from oat and from snail (*Helix aspersa*) feces. Cucumber expansins appear to assoc. with the cellulose fraction of the cell wall; they do not exhibit polysaccharide hydrolysis under a variety of assay condition and they do not cause a progressive weakening of the wall. Expansins also appear to disrupt hydrogen bonds as particularly noted with cellulose paper. These proteins have been identified in a wide variety of plant and other materials and have a variety of applications, including but not limited to agricultural and/or food applications and industrial uses such as their use in the paper industry as a catalyst for weakening the strength of paper products useful in the recycling of paper.

=> d his

(FILE 'HOME' ENTERED AT 13:55:57 ON 21 NOV 2002)

FILE 'MEDLINE, CAPLUS, EMBASE, BIOTECHDS, BIOSIS, SCISEARCH' ENTERED AT
13:56:35 ON 21 NOV 2002

L1	0 S DNA PROBE AND EXPANSIN ACTIVITY
L2	0 S DNA PROBE AND EXPANSIN
L3	1 S DNA AND PROBE AND EXPANSIN
L4	4 S EXPANSIN PROTEIN AND DNA
L5	4 DUP REM L4 (0 DUPLICATES REMOVED)
L6	0 S L5 AND PROBE

=> log y

COST IN U.S. DOLLARS

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TOTAL

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FULL ESTIMATED COST

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TOTAL

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WEST Search History

DATE: Thursday, November 21, 2002

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<i>DB=USPT; PLUR=YES; OP=ADJ</i>			
L5	3966543	14	L5
L4	3844890	21	L4
L3	4004976	6	L3
L2	5175275	7	L2
L1	5990182	1	L1

END OF SEARCH HISTORY